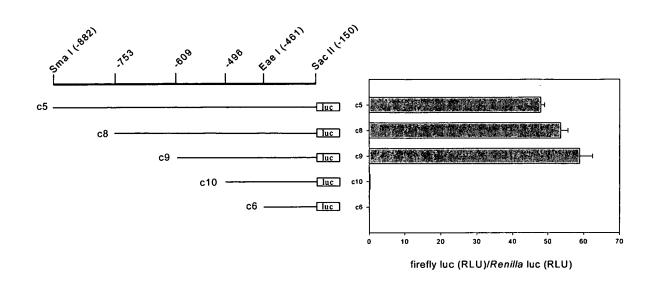
Figure 1, page 4 <VDR/RXR <SP-1 -352 GCCCTGCTGC GGTCGCCGCG AGCCTCGGCC TCTGTCTCCT CCCCCTCCCG C7 <N-Myc Hinc II Myc/Max>  $< PPAR - \alpha$ Whn> <AP-1 HNF4> CCCTTACCTC CACGCGGGAC CGCCCGCGCC AGTCAACTCC TCGCACTTTG -302  $C/EBP-\beta><E2$ < NF - YE2> TATA> A <Rel <AP-1 NF1> Ets-1> CCCCTGCTTG GCAGCGGATA AAAGGGGGCT GAGGAAATAC CGGACACGGT <RFX1 <NF-1 TATA> Whn> CACCCGTTGC CAGCTCTAGC CTTTAAATTC CCGGCTCGGG GACCTCCACG Sac II -152 CACCGCGGCT AGCGCCGACA ACCAGCTAGC GTGCAAGGCG CCGCGGCTCA -102 GCGCGTACCG GCGGGCTTCG AAACCGCAGT CCTCCGGCGA CCCCGAACTC CGCTCCGGAG CCTCAGCCCC CTGGAAAGTG ATCCCGGCAT CCGAGAGCCA (81:049IP32) a +1 Q D D V S F P M P A Η L L -2 AGATGCCGGC CCACTTGCTG CAGGACGATG TGAGTTTCCC AGCCTGGCCC (SEQ ID NO: 15)

Figur	ce 2				· • · · · · · · · · · · · · · · · · · ·
-		<arnt< th=""><th><ets-1< th=""><th></th><th>1944 5T.</th></ets-1<></th></arnt<>	<ets-1< th=""><th></th><th>1944 5T.</th></ets-1<>		1944 5T.
		< N-MyC	NF-1>	<	$PPAR-\alpha$
		• '	<sp-1< th=""><th><vdf< th=""><th>}</th></vdf<></th></sp-1<>	<vdf< th=""><th>}</th></vdf<>	}
		Whn>	< NF - kB		< HNF - 4
mSCD1	(-298)	ACCTCCACGCCTGG	CTTCCTTGGCTAG	CTATCTCTGCGCTC	CTTTA
hSCD	(-298)	ACCTCCACGCGGGA	: :: :: :: ACCGCCCGCGCCAG	: ::: ::: : GTCAACTCCTCGCAC VDR>	CTTTG
				Ets-1>	
		AP-	4>	<c-rel< td=""><td></td></c-rel<>	
			TATA>	C/EBP-β>	•
		NF-1>	GATA1>GKLF>	<hnf3-< td=""><td>·β</td></hnf3-<>	·β
mSCD1	(-253)	CCCTTTGCTGGCAG	GCCGATAAAAGGGG	GCTGAGGAAA TACT	rgaac
		:::::::::::::::::::::::::::::::::::::::		::::	: ::
hSCD	(-253)	CCCCTGCTTGGCAC	GCGGATAAAAGGGG	GCTGAGGAAATAC	CGGAC
		$RAR-\alpha1>$			
		USF>	<nf-1< td=""><td></td><td>Ets-1&gt;</td></nf-1<>		Ets-1>
			RFX-1	TATA> <ets-1< td=""><td></td></ets-1<>	
mSCD1	(-208)	ACGGTCATCCCATC	CGCCTGCTCTACCC	'TTTAAAATCCCAGC	CCAG
hSCD	(-207)	ACGGTCA-CCCGTT	::: ::::::::::::::::::::::::::::::::::	::::::::::::::::::::::::::::::::::::::	: : CTC-G
		<gata3whn></gata3whn>			
mSCD1	(-163)	GAGATCTGTGCACA	AGCCAGACCGGGCT	GAACACCCATCCC	GAGAG
	( ,	: :: :: :: ::	: : : :	::: ::: :	: :
hSCD	(-164)	GGGACCTCCACGCA	ACCGCGGCTAGCGC	CCGACAACCAGCTA	GCGTG
			_		
mSCD1	(-118)	TCAGGAGGGCAGGT	CTTCCAAGCGCAGT	TCCGCCACTCGCCT	ACAC
1 0 05	( 440)	::: : ::	: ::::: ::	::: : :	: ::
nscu	(-119)	CAAGGCGCCGCGGC	TCAGCGC-G1	ACCGGCGGGCTTCG	SAAAC
mSCD1	(-73)	CAACGGGCTCCGGA	ACCGAAGTCC	ACGCTCGATC-TC	GCAC
	, ,			: : ,:: : :::	
hSCD	(-78)	CGCAGTCCTCCGGC			
				+1	בר ישני בר
mSCD1	(-32)	TG-GGAAAGTGAGG	GGAGCAACTGACT	ATCATCATG (260	(1D/W-17)
ት C C C	( 22)		: ::: : ::	:: :::	(8) :04 OT E
nscD	(-33)	CCTGGAAAGTGATC	CCGGCATCCGAGA	GCCAAGATG (SC	4 4F/V= 14]



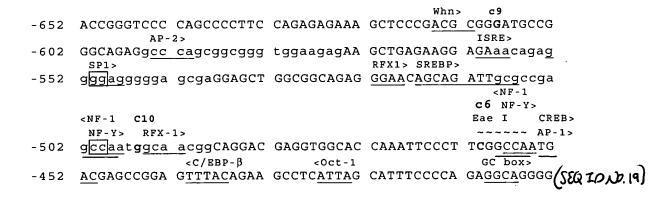


Figure 5

## Figure 8

		AP-4>
		IK2> SRE>
		IK2> RFX-1>
mSCD1	(-610)	GGGAGGAGAGCGGAGAGCTAGAGGCAGAGGGAAC <b>AGC</b>
mSCD2	(-487)	::::::::::::::::::::::::::::::::::::::
hSCD	(-552)	GGGAGG-GGGAGCGAGGGGGGGGGGGGGGGGGGGGGGGG
		CCAAT> NF-Y>
		<nf-1 <δef-1<="" td=""></nf-1>
mSCD1	(-571)	AGATTGCGCCTAGCCAATGGAAAAGGCAGGACAAGGTGG (SEQ TP NO. 20)
mSCD2	(-448)	AGATTGTGCAGAGCCAATGAGAGCAGCAGGACGAGGTGG (JEQ IP N0: 21)
heen	/_514)	AGATTECECCE CCCAATECCEAACCCCACCACCACCACCTCC (SEQ TO NO. 22)